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1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/612,809B

DATE: 02/10/2033

TIME: 15:04:09

Input Set : A:\Iowa042.app
Output Set: N:\CRF4\02102003\I612809B.raw

3 <110> APPLICANT: SHEFFIELD, VAL C.
4 AIWARI, WALLACE L.M.
5 STONE, ERWIN M.
6 NISHIMURA, DARRYL
7 PATEL, SHIVA
9 <120> TITLE OF INVENTION: THERAPEUTICS AND DIAGNOSTICS FOR CONGENITAL HEART
10 DISEASE BASED ON A NOVEL HUMAN TRANSCRIPTION FACTOR
12 <130> FILE REFERENCE: IOWA:0410USD1
14 <140> CURRENT APPLICATION NUMBER: 09/612,809B
15 <141> CURRENT FILING DATE: 2000-07-10
17 <160> NUMBER OF SEQ ID NOs: 20
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <210> LENGTH: 21884
23 <210> TYPE: cDNA
24 <210> ORGANISM: Homo sapiens
25 <210> DBSOURCE: 1

ENTER

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53 ctgtgcggcc agatatacgac agataaaat at tggcttgtg tattccatat aaaatigcag 1620
 54 tgcataattat acatccctgt gagccagatg ctgaatagat ttttccat tatttcgtc 1680
 55 ctttataaaa gggaaaataaa accagtittt aaatgtatgt atataattctt ccccaattha 1740
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 57 gggatattta atctttgaga aactattita gaaaatatgt ttgttagaaca attatitgt 1860
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 62 ggatgggggg ctgtcaatgtt gttttggaga taatacagtt tcctgtatac tgccgtct 2160
 63 atcttagagggc aacacttaaq cagtaatigc tgggtttgt tgcacaaatt tgatcattgt 2220
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 65 n/aad 2284

66 <110> SEQ ID NO: 2
 67 <111> LENGTH: 553
 68 <112> TYPE: PRT
 69 <113> ORGANISM: Homo sapiens
 70 <114> SEQUENCE: 2
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 72 1 5 10 15
 73 Pro Tyr Leu Gly G.y Glu Gln Ser Tyr Tyr Arg Ala Ala Ala Ala
 74 20 25 30
 75 Ala Gly Gly Gly Tyr Thr Ala Met Pro Ala Pro Met Ser Val Tyr Ser
 76 35 40 45
 77 His Pro Ala His Ala Glu Gln Tyr Pro Gly Gly Met Ala Arg Ala Tyr
 78 50 55 60
 79 Gly Pro Tyr Thr Pro Gln Pro Gln Pro Lys Asp Met Val Lys Pro Pro
 80 65 70 75 80
 81 Tyr Ser Tyr Ile Ala Leu Ile Thr Met Ala Ile Gln Asn Ala Pro Asp
 82 85 90 95
 83 Lys Lys Ile Thr Leu Asn Gly Ile Tyr Gln Phe Ile Met Asp Arg Phe
 84 100 105 110
 85 Pro Phe Tyr Arg Asp Asn Lys Gln Gly Trp Gln Asn Ser Ile Arg His
 86 115 120 125
 87 Asn Leu Ser Leu Asn Glu Cys Phe Val Lys Val Pro Arg Asp Asp Lys
 88 130 135 140
 89 Lys Pro Gly Lys Gly Ser Tyr Trp Thr Leu Asp Pro Asp Ser Tyr Arg
 90 145 150 155 160
 91 Met Phe Glu Asn Gly Ser Phe Leu Arg Arg Arg Arg Arg Pro Lys Lys
 92 165 170 175 180
 93 Lys Asp Ala Val Lys Asp Lys Glu Glu Lys Asp Arg Leu His Leu Lys
 94 185 190 195
 95 Glu Pro Pro Pro Pro Gly Arg Gln Pro Pro Pro Ala Pro Pro Glu Gln
 96 195 200 205
 97 Ala Asp Gly Asn Ala Pro Gly Pro Gln Pro Pro Pro Val Arg Ile Gln
 98 210 215 220
 99 Asp Ile Lys Thr Glu Asn Gly Thr Cys Pro Ser Pro Pro Gln Pro Leu
 100 225 230 235 240
 101 Ser Pro Ala Ala Ala Leu Gly Ser Gly Ser Ala Ala Ala Val Pro Lys

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253 <220> FEATURE:

254 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

255 Peptide

257 <400> SEQUENCE: 5

258 Ile Lys Asp Leu Val Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Thr

259 5 10 15

260 Met Ala Ile Gln Asn Ala Pro Glu Lys Lys Ile Thr Leu Asn Gly Ile

261 20 25 30

262 Tyr Gln Ile Met Asp Arg Phe Pro Phe Tyr Arg Glu Asn Lys Gln

263 35 40 45

264 Arg Trp Gln Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu Cys Phe

265 50 55 60

266 Val Lys Val Pro Arg Asp Asp Lys Lys Pro Gly Lys Gly Ser Tyr Trp

267 65 70 75 80

268 Thr Leu Asp Pro Asp Ser Tyr Asn Met Phe Glu Asn Gly Ser Phe Leu

269 85 90 95

270 Arg Arg Arg Arg Arg Phe Lys Lys Lys Asp

271 100 105 110

272 <410> SEQ ID NO: 5

273 <411> LENGTH: 106

274 <412> TYPE: PPT

275 <413> ORGANISM: Artificial Sequence

276 <414> FEATURE:

277 <415> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

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281 5 10 15

282 Met Ala Ile Gln Ser Ser Pro Gly Gln Arg Ala Thr Leu Ser Gly Ile

283 20 25 30

284 Tyr Arg Val Ile Met Gly Arg Phe Ala Phe Tyr Arg His Asn Arg Pro

285 35 40 45

286 Arg Trp Ile Asn Ser Ile Arg His Asn Leu Ser Leu Asn Glu Cys Phe

287 50 55 60

288 Val Lys Val Pro Arg Asp Asp Arg Lys Pro Gly Lys Gly Ser Tyr Trp

289 65 70 75 80

290 Thr Leu Asp Pro Asp Cys His Asp Met Phe Glu His Gly Ser Phe Leu

291 85 90 95

292 Arg Arg Arg Arg Arg Arg Phe Thr Arg Gln Thr

293 100 105 110

294 <410> SEQ ID NO: 7

295 <411> LENGTH: 106

296 <412> TYPE: PPT

297 <413> ORGANISM: Artificial Sequence

298 <414> FEATURE:

299 <415> OTHER INFORMATION: Description of Artificial Sequence: Synthetic

300 Peptide

301 <400> SEQUENCE: 7

302 Ala Glu Thr Pro Gin Lys Pro Pro Tyr Ser Tyr Ile Ala Leu Ile Ala

VERIFICATION SUMMARY

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